



TITLE. METHOD AND APPARATUS FOR
AUTOMATICALLY REMOVING VECTOR UNIT IN
DNA BASE SEQUENCE
INVENTORS: Kensaku IMAI, et al.
SERIAL NO.: 09/785,269
DOCKET NO.: 826,1335C

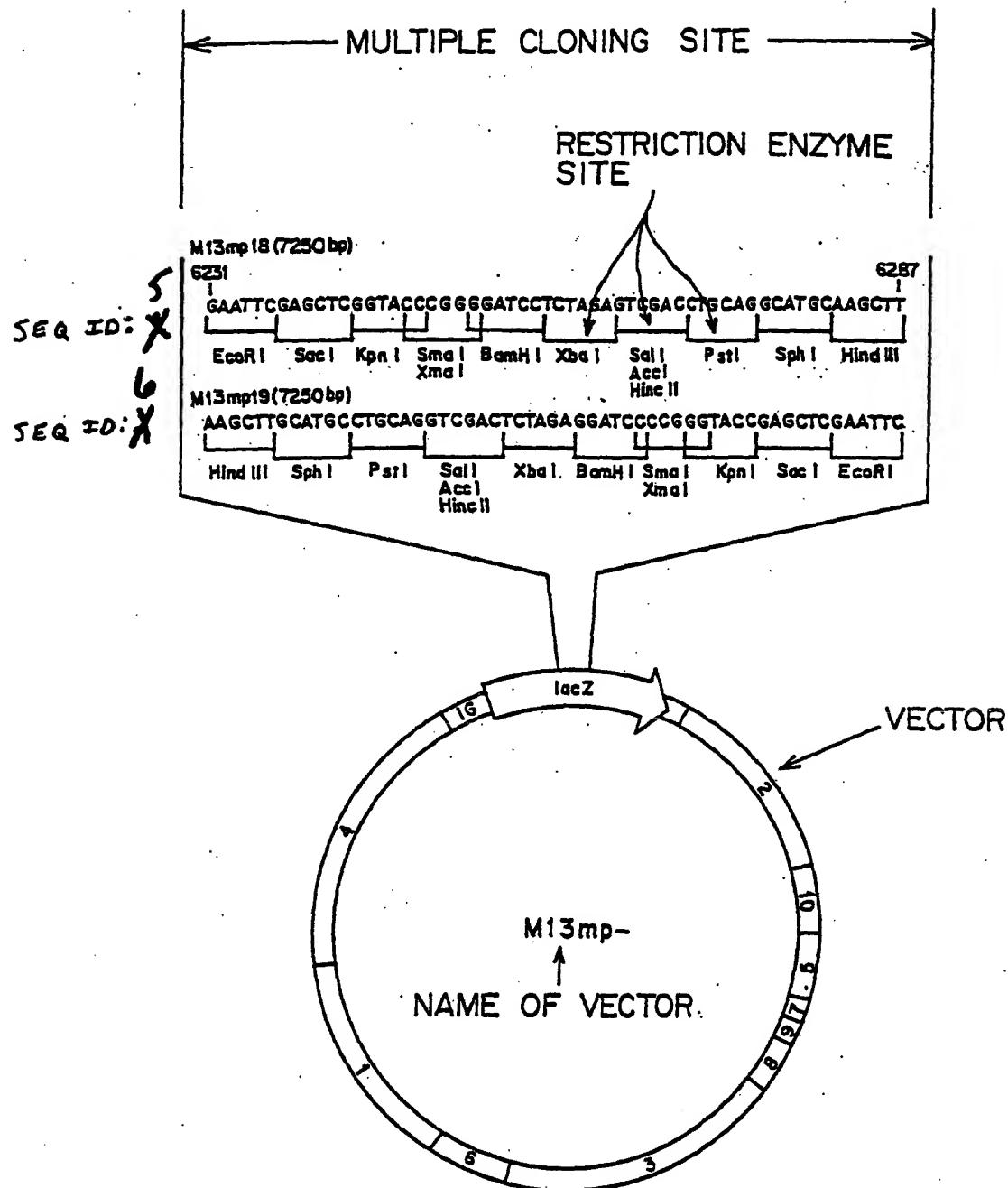
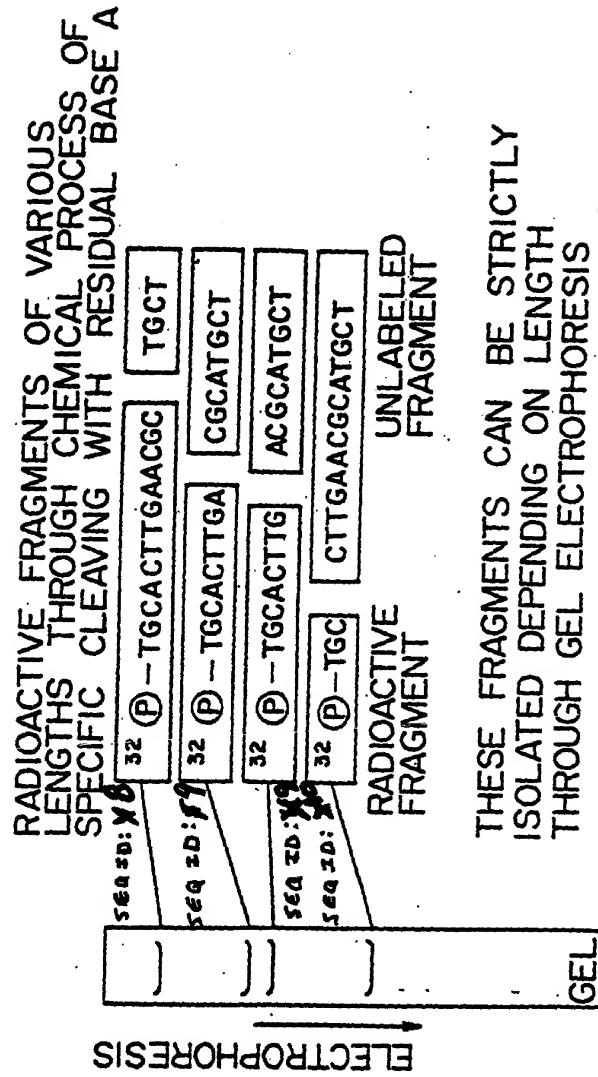


FIG. 3



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DNA FRAGMENT LABELED WITH ^{32}P AT 5' EDGE
SEQ ID: X7 ^{32}P - TGCACCTGAACCGATGCT



THESE FRAGMENTS CAN BE STRICTLY ISOLATED DEPENDING ON LENGTH THROUGH GEL ELECTROPHORESIS

FIG. 4



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VECTOR DB FORMAT

>ID

PUC18

>SEQ ID: X 11

TCGGCGTTTGGTATGACGGTAAAACCTCTGACACATGCAAGCTCCGGAGACGGTCACAGCTTGTCTGAAGCGGAT
GCCGGGAGCAGACAAGCCGTCAAGGCGCGTCAGCGGGTGTGGCGGGTGTGGGGCTGGCTTAACATGCGGCATCAGA
GCAGATTGTACTGAGAGTGCACCATATGCGGTGAAATACCGCACAGATGCGTAAGGAGAAAATACCGCATCAGGC
ATTCGCCATTCAAGGCTGCGCAACTGTTGGAAAGGGCAGTCGGTGCAGGGCTTTCGCTATTACGCCAGCTGGC
GGATGTGCTGCAAGGCATTAAAGTTGGTAACGCACGGTTTCCCAGTCAGGACGTTGAAACGACGGCAGTGCCAA
GCTTGATGCTGCAAGGCTGACTCTAGAGGATCCCGGTACCGAGCTGAATTGTAATCATGGTATAGCTGTTCT
GTGTGAAATTGTTATCCGTCACAATTCCACACAATACGAGCGGAAGCATAAAGTGAAGGCTGGGGTGCCTAATG
AGTGAAGCTAATCACAATTATGCGTTGGCTCACTGCCGCTTCCAGTCGGGAAACCTGCGCAGCTGCATTAA
GAATCGGCCAACCGCGGGAGAGGGCGTTGCGTATTGGGCGCTTCCGCTCCCGTCACTGACTCGCTGCGCTCG
GTCGTTGGCTGCGCGAGCGGTATCGCTACTCAAAGCGGTAAACGGTTATCCACAGAATCAGGGATAACGCA
AAAGAACATGTGAGCAAAGGCCAGCAAAGGCCAGGAACCGTAAACGGCGCTTGGCGTACAGGACTAAAGATA
GCCCGGCTGACGAGCATCACAAATCGACGCTCAAGTCAGGTTGGCGAACCCGACAGGACTAAAGATA
TTTCCCGCTGGAAGCTCCCTGCGCTCTCTGCGACCCGCTCCGTTACCGGATACTGTCGGCTTCTCCCTC
GGGAAGCGTGGCGCTTCTCAAAGCTACGCTGTAGGTATCTCAGTTGCGTGTAGGCTTCGCTCAAGCTGGCTGTG
TGCACGAACCCCCCGTTAGCCGACCGCTGCGCTTATCGGTAACTATGTTGAGTCAACCCGTAAGACACGAC
TTATGCCACTGGCAGCAGCACTGGTAACAGGATTAGCAGAGCAGGTATGAGGCGGTCTACAGATTGAAAGTG
GTGGCTAACTACGGCTACACTAGAAGAACAGTATTGGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCG
TTGGTAGCTTGTGATCCGGAAACAAACCCACCGCTGGTAGCGGGTTTTTTGTTGCAAGCAGCAGATTACGCG
AAAAAAGGATCTAAGAAGATCTTGTCTTGTCTTACGGGTCTGACGCTCAGTGGAAACGAAAACCTACGGTA
TTGGTCTGAGATTATCAAAAGGATCTTACCTAGATCTTAAATTAAAAAATGAAGTTAAATCAATCTAAAGTA
TATATGAGTAAACTTGGTCTGACAGTTACCAATGCTTAATCAGTGAGGCACCTATCTCAGCGATCTGCTATT
TCCATAGTTGCTGACTCCCCGCTGTAGATAACTACGATACGGGGGGCTTACCATCTGGCCCCAGTGTGCAATGAT
ACCGCGAGACCCACGCTCACCGGCTCAGATTATCAGCAATAAACCAGCCAGCCGGAAAGGGCGAGCGCAGAAGTGGTC
CTGCAACTTATCGCCCTCATCCAGTCTATTAAATTGTCGGGGAAAGCTAGAGTAAGTAGTTGCTTGGCAGTTAAG
CGCAACGTTGGTGCCTACAGGCATCGTGGTGTACGCTCGTCTGGTATGGCTTCACTCAGCTCCGGTCCCA
ACGATCAAGGCAGTTACATGATCCCCATGTTGCAAAAAAGCGGTAGCTCTCGGTCTCCGATGTTGTCAGAA
GTAAGTTGGCCGAGTGTATCACTCATGGTATGGCAGCACTGCATAATTCTTAACTGTCTATGCCATCCGTAAGATGC
TTTCTGTGACTGGTGAATCAACAAAGTCACTTCTGAGAAATAGTGTATGCGCGACCGAGTTGCTTGGCG
AATACGGGATAATACCGGCCACATAGCAGAACTTTAAAGTGTCTATCATTGGAAAACGTTCTGGGGCGAAA
ACTCTCAAGGATCTTACCGCTGTTGAGATCCAGTTGATGTAACCCACTCGTCACCCAACTGATCTCAGCAT
ACCAGCGTTCTGGGTGAGCAAAACAGGAAGGCAAATGCCGAAAAAGGGATAAGGGCGACACGGAAATGTTGAAT
ACTCATACTCTCCTTTCAATTATTGAAGCATTTACGGGTATTGTCATGAGGGATAACATATTGAATGTA
TTAGAAAAATAACAAATAGGGTCCGCGACATTCCCGAAAAGTGCACCTGACGTCTAAGAAACCA
ATGACATTAACCTATAAAAGGCATACGAGGCCCTTCGTC

>MULTI

399..450

FIG. 9



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(* INDICATES MULTIPLE CLONING SITE)

SEQ ID: 8 GTGCCAAGCTTGCATGCCCTGCAGGTCGACTCTAGAGGATCCCCGGTACCGAGCTCGAAATTCTGTAAT
SEQ ID: 9 ¹² AAGCTT¹³ → HIND III
SEQ ID: 10 ¹⁴ GCATGC¹⁵ → SPH I
SEQ ID: 11 ¹⁶ CTGCAG¹⁷ → PST I
SEQ ID: 12 ¹⁸ GTCGAC¹⁹ → SAL I, ACC I, HINC II
SEQ ID: 13 ²⁰ TCTAGA²¹ → XBA I
SEQ ID: 14 ²² GGATCC²³ → BAMH I
SEQ ID: 15 ²⁴ CCCGGG²⁵ → SMA I, XMA I
SEQ ID: 16 ²⁶ GGTACC²⁷ → KPN I
SEQ ID: 17 ²⁸ GAGCTC²⁹ → SAC I
SEQ ID: 18 ³⁰ GAATTTC³¹ → BCG I

FIG. 10

WHEN HIND III IS SPECIFIED ON VECTOR 5' SIDE
XBA I IS SPECIFIED ON VECTOR 3' SIDE, HIND III IS
SPECIFIED ON OBJECT DNA 5' SIDE, AND XBA I IS
SPECIFIED ON OBJECT DNA 3' SIDE

(**** INDICATES RESIDUAL MULTIPLECLONING SITE
(+ + + + INDICATES AN OBJECT DNA FRAGMENT

(SEQUENCE
ID NO. 4)

GTGCCAAGCTT ++++++ TCTAGAGGATCCCCGGGTACCGAGCTCGAATTCTGAAT
AAGCTT TCTAGA
↑ ↑

5' SIDE RETRIEVAL KEY
(IN THIS EXAMPLE,
HIND III SITE)

***** (SEQUENCE
ID NO. 23)
CTAGAGGATCCCCGGTACCGAGCTCGAATTCTGTAAT
CTAGA
↑

9 SIDE RETRIEVAL KEY
(IN THIS EXAMPLE, XBA I SITE)

FIG. 17